

RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.**

Application Serial Number: 10/567,731
Source: TFWP
Date Processed by STIC: 01/11/2007

ENTERED



IFWP

RAW SEQUENCE LISTING

DATE: 01/11/2007

PATENT APPLICATION: US/10/567,731

TIME: 10:56:49

Input Set : A:\Sequence Listing.txt

Output Set: N:\CRF4\01112007\J567731.raw

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3 <110> APPLICANT: TAKARA BIO INC.
5 <120> TITLE OF INVENTION: Method of decomposing dsRNA and synthesizing RNA
7 <130> FILE REFERENCE: 664674
C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/567,731
C--> 9 <141> CURRENT FILING DATE: 2006-02-10
9 <150> PRIOR APPLICATION NUMBER: JP 2003-293553
10 <151> PRIOR FILING DATE: 2003-08-14
12 <150> PRIOR APPLICATION NUMBER: JP 2003-342126
13 <151> PRIOR FILING DATE: 2003-09-30
15 <150> PRIOR APPLICATION NUMBER: JP 2003-409639
16 <151> PRIOR FILING DATE: 2003-12-08
18 <150> PRIOR APPLICATION NUMBER: JP 2004-086129
19 <151> PRIOR FILING DATE: 2004-03-24
21 <160> NUMBER OF SEQ ID NOS: 27
23 <170> SOFTWARE: PatentIn version 3.1
25 <210> SEQ ID NO: 1
26 <211> LENGTH: 1924
27 <212> TYPE: PRT
28 <213> ORGANISM: Homo sapiens
30 <400> SEQUENCE: 1
32 Met Lys Ser Pro Ala Leu Gln Pro Leu Ser Met Ala Gly Leu Gln Leu
33 1 5 10 15
36 Met Thr Pro Ala Ser Ser Pro Met Gly Pro Phe Phe Gly Leu Pro Trp
37 20 25 30
40 Gln Gln Glu Ala Ile His Asp Asn Ile Tyr Thr Pro Arg Lys Tyr Gln
41 35 40 45
44 Val Glu Leu Leu Glu Ala Ala Leu Asp His Asn Thr Ile Val Cys Leu
45 50 55 60
48 Asn Thr Gly Ser Gly Lys Thr Phe Ile Ala Ser Thr Thr Leu Leu Lys
49 65 70 75 80
52 Ser Cys Leu Tyr Leu Asp Leu Gly Glu Thr Ser Ala Arg Asn Gly Lys
53 85 90 95
56 Arg Thr Val Phe Leu Val Asn Ser Ala Asn Gln Val Ala Gln Gln Val
57 100 105 110
60 Ser Ala Val Arg Thr His Ser Asp Leu Lys Val Gly Glu Tyr Ser Asn
61 115 120 125
64 Leu Glu Val Asn Ala Ser Trp Thr Lys Glu Arg Trp Asn Gln Glu Phe
65 130 135 140
68 Thr Lys His Gln Val Leu Ile Met Thr Cys Tyr Val Ala Leu Asn Val
69 145 150 155 160
72 Leu Lys Asn Gly Tyr Leu Ser Leu Ser Asp Ile Asn Leu Leu Val Phe
73 165 170 175
76 Asp Glu Cys His Leu Ala Ile Leu Asp His Pro Tyr Arg Glu Phe Met

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77      180      185      190
80 Lys Leu Cys Glu Ile Cys Pro Ser Cys Pro Arg Ile Leu Gly Leu Thr
81      195      200      205
84 Ala Ser Ile Leu Asn Gly Lys Trp Asp Pro Glu Asp Leu Glu Glu Lys
85      210      215      220
88 Phe Gln Lys Leu Glu Lys Ile Leu Lys Ser Asn Ala Glu Thr Ala Thr
89 225      230      235      240
92 Asp Leu Val Val Leu Asp Arg Tyr Thr Ser Gln Pro Cys Glu Ile Val
93      245      250      255
96 Val Asp Cys Gly Pro Phe Thr Asp Arg Ser Gly Leu Tyr Glu Arg Leu
97      260      265      270
100 Leu Met Glu Leu Glu Glu Ala Leu Asn Phe Ile Asn Asp Cys Asn Ile
101      275      280      285
104 Ser Val His Ser Lys Glu Arg Asp Ser Thr Leu Ile Ser Lys Gln Ile
105      290      295      300
108 Leu Ser Asp Cys Arg Ala Val Leu Val Val Leu Gly Pro Trp Cys Ala
109 305      310      315      320
112 Asp Lys Val Ala Gly Met Met Val Arg Glu Leu Gln Lys Tyr Ile Lys
113      325      330      335
116 His Glu Gln Glu Glu Leu His Arg Lys Phe Leu Leu Phe Thr Asp Thr
117      340      345      350
120 Phe Leu Arg Lys Ile His Ala Leu Cys Glu Glu His Phe Ser Pro Ala
121      355      360      365
124 Ser Leu Asp Leu Lys Phe Val Thr Pro Lys Val Ile Lys Leu Leu Glu
125      370      375      380
128 Ile Leu Arg Lys Tyr Lys Pro Tyr Glu Arg His Ser Phe Glu Ser Val
129 385      390      395      400
132 Glu Trp Tyr Asn Asn Arg Asn Gln Asp Asn Tyr Val Ser Trp Ser Asp
133      405      410      415
136 Ser Glu Asp Asp Asp Glu Asp Glu Glu Ile Glu Glu Lys Glu Lys Pro
137      420      425      430
140 Glu Thr Asn Phe Pro Ser Pro Phe Thr Asn Ile Leu Cys Gly Ile Ile
141      435      440      445
144 Phe Val Glu Arg Arg Tyr Thr Ala Val Val Leu Asn Arg Leu Ile Lys
145      450      455      460
148 Glu Ala Gly Lys Gln Asp Pro Glu Leu Ala Tyr Ile Ser Ser Asn Phe
149 465      470      475      480
152 Ile Thr Gly His Gly Ile Gly Lys Asn Gln Pro Arg Asn Asn Thr Met
153      485      490      495
156 Glu Ala Glu Phe Arg Lys Gln Glu Glu Val Leu Arg Lys Phe Arg Ala
157      500      505      510
160 His Glu Thr Asn Leu Leu Ile Ala Thr Ser Ile Val Glu Glu Gly Val
161      515      520      525
164 Asp Ile Pro Lys Cys Asn Leu Val Val Arg Phe Asp Leu Pro Thr Glu
165      530      535      540
168 Tyr Arg Ser Tyr Val Gln Ser Lys Gly Arg Ala Arg Ala Pro Ile Ser
169 545      550      555      560
172 Asn Tyr Ile Met Leu Ala Asp Thr Asp Lys Ile Lys Ser Phe Glu Glu
173      565      570      575

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176 Asp Leu Lys Thr Tyr Lys Ala Ile Glu Lys Ile Leu Arg Asn Lys Cys
177          580          585          590
180 Ser Lys Ser Val Asp Thr Gly Glu Thr Asp Ile Asp Pro Val Met Asp
181          595          600          605
184 Asp Asp His Val Phe Pro Pro Tyr Val Leu Arg Pro Asp Asp Gly Gly
185          610          615          620
188 Pro Arg Val Thr Ile Asn Thr Ala Ile Gly His Ile Asn Arg Tyr Cys
189 625          630          635          640
192 Ala Arg Leu Pro Ser Asp Pro Phe Thr His Leu Ala Pro Lys Cys Arg
193          645          650          655
196 Thr Arg Glu Leu Pro Asp Gly Thr Phe Tyr Ser Thr Leu Tyr Leu Pro
197          660          665          670
200 Ile Asn Ser Pro Leu Arg Ala Ser Ile Val Gly Pro Pro Met Ser Cys
201          675          680          685
204 Val Arg Leu Ala Glu Arg Val Val Ala Leu Ile Cys Cys Glu Lys Leu
205          690          695          700
208 His Lys Ile Gly Glu Leu Asp Asp His Leu Met Pro Val Gly Lys Glu
209 705          710          715          720
212 Thr Val Lys Tyr Glu Glu Glu Leu Asp Leu His Asp Glu Glu Glu Thr
213          725          730          735
216 Ser Val Pro Gly Arg Pro Gly Ser Thr Lys Arg Arg Gln Cys Tyr Pro
217          740          745          750
220 Lys Ala Ile Pro Glu Cys Leu Arg Asp Ser Tyr Pro Arg Pro Asp Gln
221          755          760          765
224 Pro Cys Tyr Leu Tyr Val Ile Gly Met Val Leu Thr Thr Pro Leu Pro
225          770          775          780
228 Asp Glu Leu Asn Phe Arg Arg Arg Lys Leu Tyr Pro Pro Glu Asp Thr
229 785          790          795          800
232 Thr Arg Cys Phe Gly Ile Leu Thr Ala Lys Pro Ile Pro Gln Ile Pro
233          805          810          815
236 His Phe Pro Val Tyr Thr Arg Ser Gly Glu Val Thr Ile Ser Ile Glu
237          820          825          830
240 Leu Lys Lys Ser Gly Phe Met Leu Ser Leu Gln Met Leu Glu Leu Ile
241          835          840          845
244 Thr Arg Leu His Gln Tyr Ile Phe Ser His Ile Leu Arg Leu Glu Lys
245          850          855          860
248 Pro Ala Leu Glu Phe Lys Pro Thr Asp Ala Asp Ser Ala Tyr Cys Val
249 865          870          875          880
252 Leu Pro Leu Asn Val Val Asn Asp Ser Ser Thr Leu Asp Ile Asp Phe
253          885          890          895
256 Lys Phe Met Glu Asp Ile Glu Lys Ser Glu Ala Arg Ile Gly Ile Pro
257          900          905          910
260 Ser Thr Lys Tyr Thr Lys Glu Thr Pro Phe Val Phe Lys Leu Glu Asp
261          915          920          925
264 Tyr Gln Asp Ala Val Ile Ile Pro Arg Tyr Arg Asn Phe Asp Gln Pro
265          930          935          940
268 His Arg Phe Tyr Val Ala Asp Val Tyr Thr Asp Leu Thr Pro Leu Ser
269 945          950          955          960
272 Lys Phe Pro Ser Pro Glu Tyr Glu Thr Phe Ala Glu Tyr Tyr Lys Thr

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273		965		970		975
276	Lys	Tyr	Asn	Leu	Asp	Leu
277			980		985	
280	Asp	His	Thr	Ser	Ser	Arg
281			995		1000	
284	Gln	Lys	Gly	Lys	Ala	Leu
285		1010			1015	
288	Ala	Lys	Trp	Glu	Ser	Leu
289		1025			1030	
292	Leu	Cys	Ala	Ile	His	Pro
293		1040			1045	
296	Val	Cys	Leu	Pro	Ser	Ile
297		1055			1060	
300	Ala	Glu	Glu	Leu	Arg	Ala
301		1070			1075	
304	Val	Arg	Ser	Leu	Pro	Ala
305		1085			1090	
308	Gly	Trp	Lys	Lys	Ser	Ile
309		1100			1105	
312	Asn	Ser	Ser	Ser	Ala	Glu
313		1115			1120	
316	Ile	Val	Pro	Glu	Asn	Ala
317		1130			1135	
320	Ser	Leu	Glu	Asn	His	Asp
321		1145			1150	
324	Leu	Ser	Glu	Ser	Pro	Gly
325		1160			1165	
328	Leu	Thr	Ala	Ile	Asn	Gly
329		1175			1180	
332	Gly	Ser	Tyr	Asp	Leu	Ala
333		1190			1195	
336	Leu	Asn	Tyr	Tyr	Lys	Gln
337		1205			1210	
340	Tyr	Ser	Ile	Gln	Asn	Leu
341		1220			1225	
344	Ser	Asp	Glu	Cys	Thr	Leu
345		1235			1240	
348	Ala	Asn	Lys	Ser	Thr	Ser
349		1250			1255	
352	Pro	Gly	Thr	Thr	Asp	Thr
353		1265			1270	
356	Ser	Glu	Gln	Ser	Pro	Ser
357		1280			1285	
360	Pro	Asn	Pro	Gly	Leu	Ile
361		1295			1300	
364	Ser	Asp	Gly	Phe	Asn	Leu
365		1310			1315	
368	Phe	Leu	Lys	His	Ala	Ile
369		1325			1330	

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372 Asp Ala His Glu Gly Arg Leu Ser Tyr Met Arg Ser Lys Lys Val
373      1340      1345      1350
376 Ser Asn Cys Asn Leu Tyr Arg Leu Gly Lys Lys Lys Gly Leu Pro
377      1355      1360      1365
380 Ser Arg Met Val Val Ser Ile Phe Asp Pro Pro Val Asn Trp Leu
381      1370      1375      1380
384 Pro Pro Gly Tyr Val Val Asn Gln Asp Lys Ser Asn Thr Asp Lys
385      1385      1390      1395
388 Trp Glu Lys Asp Glu Met Thr Lys Asp Cys Met Leu Ala Asn Gly
389      1400      1405      1410
392 Lys Leu Asp Glu Asp Tyr Glu Glu Glu Asp Glu Glu Glu Glu Ser
393      1415      1420      1425
396 Leu Met Trp Arg Ala Pro Lys Glu Glu Ala Asp Tyr Glu Asp Asp
397      1430      1435      1440
400 Phe Leu Glu Tyr Asp Gln Glu His Ile Arg Phe Ile Asp Asn Met
401      1445      1450      1455
404 Leu Met Gly Ser Gly Ala Phe Val Lys Lys Ile Ser Leu Ser Pro
405      1460      1465      1470
408 Phe Ser Thr Thr Asp Ser Ala Tyr Glu Trp Lys Met Pro Lys Lys
409      1475      1480      1485
412 Ser Ser Leu Gly Ser Met Pro Phe Ser Ser Asp Phe Glu Asp Phe
413      1490      1495      1500
416 Asp Tyr Ser Ser Trp Asp Ala Met Cys Tyr Leu Asp Pro Ser Lys
417      1505      1510      1515
420 Ala Val Glu Glu Asp Asp Phe Val Val Gly Phe Trp Asn Pro Ser
421      1520      1525      1530
424 Glu Glu Asn Cys Gly Val Asp Thr Gly Lys Gln Ser Ile Ser Tyr
425      1535      1540      1545
428 Asp Leu His Thr Glu Gln Cys Ile Ala Asp Lys Ser Ile Ala Asp
429      1550      1555      1560
432 Cys Val Glu Ala Leu Leu Gly Cys Tyr Leu Thr Ser Cys Gly Glu
433      1565      1570      1575
436 Arg Ala Ala Gln Leu Phe Leu Cys Ser Leu Gly Leu Lys Val Leu
437      1580      1585      1590
440 Pro Val Ile Lys Arg Thr Asp Arg Glu Lys Ala Leu Cys Pro Thr
441      1595      1600      1605
444 Arg Glu Asn Phe Asn Ser Gln Gln Lys Asn Leu Ser Val Ser Cys
445      1610      1615      1620
448 Ala Ala Ala Ser Val Ala Ser Ser Arg Ser Ser Val Leu Lys Asp
449      1625      1630      1635
452 Ser Glu Tyr Gly Cys Leu Lys Ile Pro Pro Arg Cys Met Phe Asp
453      1640      1645      1650
456 His Pro Asp Ala Asp Lys Thr Leu Asn His Leu Ile Ser Gly Phe
457      1655      1660      1665
460 Glu Asn Phe Glu Lys Lys Ile Asn Tyr Arg Phe Lys Asn Lys Ala
461      1670      1675      1680
464 Tyr Leu Leu Gln Ala Phe Thr His Ala Ser Tyr His Tyr Asn Thr
465      1685      1690      1695
468 Ile Thr Asp Cys Tyr Gln Arg Leu Glu Phe Leu Gly Asp Ala Ile

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RAW SEQUENCE LISTING ERROR SUMMARY DATE: 01/11/2007
PATENT APPLICATION: US/10/567,731 TIME: 10:56:50

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Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:7; Line(s) 1013

Seq#:8; Line(s) 1025

Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete,
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:14,15,16,17,18,19,20,21,22,23,24,25,26,27

VERIFICATION SUMMARY

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Input Set : A:\Sequence Listing.txt

Output Set: N:\CRF4\01112007\J567731.raw

L:9 M:270 C: Current Application Number differs, Replaced Current Application No
L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date